

(SEQUENCE LISTING)

SEQ ID NO: 1

Length of sequence: 18

Type of sequence: Nucleic acid

5 Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Corresponding to from +176 to +193 of

10 SEQ ID NO: 9

Sequence:

AGAAAGTCTT CACTCTGC

SEQ ID NO: 2

15 Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

20 Antisense: YES

Feature of sequence: Corresponding to from +181 to +198 of

SEQ ID NO: 9

Sequence:

TTGAAAGAAA GTCTTCAC

25

SEQ ID NO: 3

Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

5 Antisense: YES

Feature of sequence: Corresponding to from +367 to +384 of

SEQ ID NO:9

Sequence:

GGTCTTCAGG TTCTCCCC

10

SEQ ID NO: 4

Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

15 Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Corresponding to from +637 to +654 of

SEQ ID NO:9

20 Sequence:

CTGGGTCAGC TATCCCAG

SEQ ID NO: 5

Length of sequence: 18

25 Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Corresponding to from +915 to +932 of
SEQ ID NO:9

5 Sequence:

GCTTGGAATG GAAGCTTC

SEQ ID NO: 6

Length of sequence: 18

10 Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

Kind of sequence: Another nucleic acid Synthesized DNA

Antisense: YES

15 Feature of sequence: Corresponding to from +1246 to +1263 of
SEQ ID NO:9

Sequence:

GGCTGGTTAG GAACTCCT

20 SEQ ID NO: 7

Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

25 Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Corresponding to from +1249 to +1266 of

SEQ ID NO:9

Sequence:

CCAGGCTGGT TAGGAACT

5 SEQ ID NO: 8

Length of sequence: 18

Type of sequence: Nucleic acid

Number of strand: Single-strand

Topology: Straight

10 Kind of sequence: Another nucleic acid Artificial DNA

Antisense: YES

Feature of sequence: Mouse IL-10 protein gene

Sequence:

AGGTCCTGGA GTCCAGCA

15

SEQ ID NO: 9

Length of sequence: 1601

Type of sequence: Nucleic acid

Number of strand: Single-strand

20 Topology: Straight

Kind of sequence: cDNA

Antisense: NO

Feature of sequence: cDNA of Human IL-10 protein

Sequence:

25 AAACCACAAG ACAGACTTGC AAAAGAAGGC ATGCACAGCT 40
CAGCACTGCT CTGTTGCCTG GTCCTCCTGA CTGGGGTGAG 80
GGCCAGCCCA GGCCAGGGCA CCCAGTCTGA GAACAGCTGC 120

	ACCCACTTCC	CAGGCAACCT	GCCTAACATG	CTTCGAGATC	160
	TCCGAGATGC	CTTCAGCAGA	GTGAAGACTT	TCTTTCAAAT	200
	GAAGGATCAG	CTGGACAACCT	TGTTGTTAAA	GGAGTCCTTG	240
	CTGGAGGACT	TTAAGGGTTA	CCTGGGTTGC	CAAGCCTTGT	280
5	CTGAGATGAT	CCAGTTTTAC	CTGGAGGAGG	TGATGCCCCA	320
	AGCTGAGAAC	CAAGACCCAG	ACATCAAGGC	GCATGTGAAC	360
	TCCCTGGGGG	AGAACCTGAA	GACCCTCAGG	CTGAGGCTAC	400
	GGCGCTGTCA	TCGATTTCTT	CCCTGTGAAA	ACAAGAGCAA	440
	GGCCGTGGAG	CAGGTGAAGA	ATGCCTTTAA	TAAGCTCCAA	480
10	GAGAAAGGCA	TCTACAAAGC	CATGAGTGAG	TTTGACATCT	520
	TCATCAACTA	CATAGAAGCC	TACATGACAA	TGAAGATACG	560
	AAACTGAGAC	ATCAGGGTGG	CGACTCTATA	GACTCTAGGA	600
	CATAAATTAG	AGGTCTCCAA	AATCGGATCT	GGGGCTCTGG	640
	GATAGCTGAC	CCAGCCCCTT	GAGAAACCTT	ATTGTACCTC	680
15	TCTTATAGAA	TATTTATTAC	CTCTGATACC	TCAACCCCCA	720
	TTTCTATTTA	TTTACTGAGC	TTCTCTGTGA	ACGATTTAGA	760
	AAGAAGCCCA	ATATTATAAT	TTTTTTCAAT	ATTTATTATT	800
	TTCACCTGTT	TTTAAGCTGT	TTCCATAGGG	TGACACACTA	840
	TGGTATTTGA	GTGTTTTAAG	ATAAATTATA	AGTTACATAA	880
20	GGGAGGAAAA	AAAATGTTCT	TTGGGGAGCC	AACAGAAGCT	920
	TCCATTCCAA	GCCTGACCAC	GCTTTCTAGC	TGTTGAGCTG	960
	TTTTCCCTGA	CCTCCCTCTA	ATTTATCTTG	TCTCTGGGCT	1000
	TGGGGCTTCC	TAAGTGCTAC	AAATACTCTT	AGGAAGAGAA	1040
	ACCAGGGAGC	CCCTTTGATG	ATTAATTCAC	CTTCCAGTGT	1080
25	CTCGGAGGGA	TTCCCCTAAC	CTCATTCCCC	AACCACTTCA	1120
	TTCTTGAAAG	CTGTGGCCAG	CTTGTTATTT	ATAACAACCT	1160
	AAATTTGGTT	CTAGGCCGGG	CGCGGTGGCT	CACGCCTGTA	1200

ATCCCAGCAC TTTGGGAGGC TGAGGCGGGT GGATCACTTG 1240
AGGTCAGGAG TTCCTAACCA GCCTGGTCAA CATGGTGAAA 1280
CCCCGTCTCT ACTAAAAATA CAAAATTAG CCGGGCATGG 1320
TGGCGCGCAC CTGTAATCCC AGCTACTTGG GAGGCTGAGG 1360
5 CAAGAGAATT GCTTGAACCC AGGAGATGGA AGTTGCAGTG 1400
AGCTGATATC ATGCCCCTGT ACTCCAGCCT GGGTGACAGA 1440
GCAAGACTCT GTCTCAAAAA AATAAAAATA AAAATAAATT 1480
TGGTTCTAAT AGAACTCAGT TTAACTAGA ATTTATTCAA 1520
TTCCTCTGGG AATGTTACAT TGTTTGTCTG TCTTCATAGC 1560
10 AGATTTTAAT TTTGAATAAA TAAATGTATC TTATTCACAT 1600
C 1601